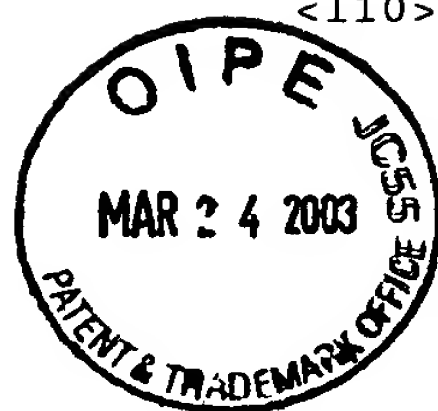


SEQUENCE LISTING



<110> Stephen M. Allen
 Gary M. Fader
 Saverio Carl Falco
 Anthony J. Kinney
 Jonathan E. Lightner
 Guo-Hua Miao
 J. Antoni Rafalski
 Catherine J. Thorpe

<120> Plant Cellulose Synthases

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RECEIVED

MAR 26 2003

TECH CENTER 1600/2900

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<212> PRT
<213> Zea mays

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Ile Asn Leu Ser Asp Arg Leu His Gln Val Leu Arg Trp Ala Leu Gly
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Ser Val Glu Ile Phe Met Ser Arg His Cys Pro Leu Trp Tyr Ala Tyr
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<212> DNA

<213> Zea mays

<400> 7

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3786

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<211> 1165

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Glu	Arg	Ser	Pro	Arg	Pro	Gly	Asp	Gln	Arg	Arg	Gly	Gly	Leu	Arg	Ala
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Phe	Arg	Cys	Ala	Ala	Ala	Ala	Gly	Phe	Val	Arg	Glu	Arg	Asp	Pro	Ala
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Gly	Arg	Gly	Gly	Gly	Pro	Glu	Met	Glu	Ala	Ser	Ala	Gly	Leu	Val	Ala
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Gly	Ser	His	Asn	Arg	Asn	Glu	Leu	Val	Val	Ile	Arg	Arg	Asp	Arg	Glu
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Ser	Gly	Ala	Ala	Gly	Gly	Gly	Ala	Ala	Arg	Arg	Ala	Glu	Ala	Pro	Cys
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Gln	Ile	Cys	Gly	Asp	Glu	Val	Gly	Val	Gly	Phe	Asp	Gly	Glu	Pro	Phe
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Val	Ala	Cys	Asn	Glu	Cys	Ala	Phe	Pro	Val	Cys	Arg	Ala	Cys	Tyr	Glu
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Tyr	Glu	Arg	Arg	Glu	Gly	Ser	Gln	Ala	Cys	Pro	Gln	Cys	Arg	Thr	Arg
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Tyr	Lys	Arg	Leu	Lys	Gly	Cys	Pro	Arg	Val	Ala	Gly	Asp	Glu	Glu	Glu
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Asp	Gly	Val	Asp	Asp	Leu	Glu	Gly	Glu	Phe	Gly	Leu	Gln	Asp	Gly	Ala
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Ala	His	Glu	Asp	Asp	Pro	Gln	Tyr	Val	Ala	Glu	Ser	Met	Leu	Arg	Ala
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Gln	Met	Ser	Tyr	Gly	Arg	Gly	Gly	Asp	Ala	His	Pro	Gly	Phe	Ser	Pro
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Pro	Lys	Trp	Leu	Pro	Ile	Glu	Arg	Glu	Thr	Tyr	Leu	Asp	Arg	Leu	Ser	
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	995			1000		1005
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 <212> PRT
 <213> Zea mays

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          20              25              30

Glu Gln Asn Gly Gln Val Cys Gln Ile Cys Gly Asp Asp Val Gly Leu
          35              40              45

Ala Pro Gly Gly Asp Pro Phe Val Ala Cys Asn Glu Cys Ala Phe Pro
          50              55              60

Val Cys Arg Asp Cys Tyr Glu Tyr Glu Arg Arg Glu Gly Thr Gln Asn
  65              70              75              80

Cys Pro Gln Cys Lys Thr Arg Tyr Lys Arg Leu Lys Gly Cys Gln Arg
          85              90              95

Val Thr Gly Asp Glu Glu Glu Asp Gly Val Asp Asp Leu Asp Asn Glu
          100             105             110

Phe Asn Trp Asp Gly His Asp Ser Gln Ser Val Ala Glu Ser Met Leu
          115             120             125

Tyr Gly His Met Ser Tyr Gly Arg Gly Gly Asp Pro Asn Gly Ala Pro
          130             135             140

Gln Ala Phe Gln Leu Asn Pro Asn Val Pro Leu Leu Thr Asn Gly Gln
          145             150             155             160

Met Val Asp Asp Ile Pro Pro Glu Gln His Ala Leu Val Pro Ser Phe
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Met Gly Gly Gly Gly Lys Arg Ile His Pro Leu Pro Tyr Ala Asp Pro
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Ser	Leu	Pro	Val	Gln	Pro	Arg	Ser	Met	Asp	Pro	Ser	Lys	Asp	Leu	Ala	195	200	205	
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Lys	Gln	Arg	Gln	Glu	Arg	Met	His	Gln	Thr	Gly	Asn	Asp	Gly	Gly	Gly	225	230	235	240
Asp	Asp	Gly	Asp	Asp	Ala	Asp	Leu	Pro	Leu	Met	Asp	Glu	Ala	Arg	Gln	245	250	255	
Gln	Leu	Ser	Arg	Lys	Ile	Pro	Leu	Pro	Ser	Ser	Gln	Ile	Asn	Pro	Tyr	260	265	270	
Arg	Met	Ile	Ile	Ile	Ile	Arg	Leu	Val	Val	Leu	Gly	Phe	Phe	Phe	His	275	280	285	
Tyr	Arg	Val	Met	His	Pro	Val	Asn	Asp	Ala	Phe	Ala	Leu	Trp	Leu	Ile	290	295	300	
Ser	Val	Ile	Cys	Glu	Ile	Trp	Phe	Ala	Met	Ser	Trp	Ile	Leu	Asp	Gln	305	310	315	320
Phe	Pro	Lys	Trp	Phe	Pro	Ile	Glu	Arg	Glu	Thr	Tyr	Leu	Asp	Arg	Leu	325	330	335	
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Lys	Val	Ser	Cys	Tyr	Val	Ser	Asp	Asp	Gly	Ala	Ala	Met	Leu	Thr	Phe	385	390	395	400
Glu	Ala	Leu	Ser	Glu	Thr	Ser	Glu	Phe	Ala	Lys	Lys	Trp	Val	Pro	Phe	405	410	415	
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Gln	Lys	Ile	Asp	Tyr	Leu	Lys	Asp	Lys	Val	Ala	Ala	Asn	Phe	Val	Arg	435	440	445	
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Asn	Ala	Leu	Val	Ala	Lys	Ala	Gln	Lys	Val	Pro	Glu	Glu	Gly	Trp	Thr	465	470	475	480
Met	Gln	Asp	Gly	Thr	Pro	Trp	Pro	Gly	Asn	Asn	Val	Arg	Asp	His	Pro	485	490	495	

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Gly	Asn	Glu	Leu	Pro	Arg	Leu	Val	Tyr	Val	Ser	Arg	Glu	Lys	Arg	Pro	515	520	525
Gly	Tyr	Asn	His	His	Lys	Lys	Ala	Gly	Ala	Met	Asn	Ala	Leu	Val	Arg	530	535	540
Val	Ser	Ala	Val	Leu	Thr	Asn	Ala	Pro	Tyr	Leu	Leu	Asn	Leu	Asp	Cys	545	550	555
Asp	His	Tyr	Ile	Asn	Asn	Ser	Lys	Ala	Ile	Lys	Glu	Ala	Met	Cys	Phe	565	570	575
Met	Met	Asp	Pro	Leu	Leu	Gly	Lys	Lys	Val	Cys	Tyr	Val	Gln	Phe	Pro	580	585	590
Gln	Arg	Phe	Asp	Gly	Ile	Asp	Arg	His	Asp	Arg	Tyr	Ala	Asn	Arg	Asn	595	600	605
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Lys	Thr	Asp	Trp	Gly	Lys	Glu	Ile	Gly	Trp	Ile	Tyr	Gly	Ser	Val	Thr	770	775	780
Glu	Asp	Ile	Leu	Thr	Gly	Phe	Lys	Met	His	Cys	His	Gly	Trp	Arg	Ser	785	790	795
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Ile Tyr Cys Ile Pro Lys Arg Val Ala Phe Lys Gly Ser Ala Pro Leu
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 Ile Glu Ile Phe Phe Ser Asn His Cys Pro Leu Trp Tyr Gly Tyr Gly
 835 840 845
 Gly Gly Leu Lys Phe Leu Glu Arg Phe Ser Tyr Ile Asn Ser Ile Val
 850 855 860
 Tyr Pro Trp Thr Ser Ile Pro Leu Leu Ala Tyr Cys Thr Leu Pro Ala
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<213> Oryza sativa

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<213> Oryza sativa

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Ala Ala Leu Trp Val Ser Phe Cys Arg Lys His Gly Val Glu Pro Arg
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Asn Leu Glu Ala Tyr Phe Asn Ala Gly Glu Gly Gly Gly Gly Lys Ala
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Lys Val Val Ala Arg Gly Ser Tyr Arg Gly Met Ala Trp Pro Glu Leu
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Val Arg Asp Arg Arg Arg Val Arg Arg Glu Tyr Glu Glu Met Arg Leu
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Arg Ile Asp Ala Leu Gln Ala Ala Asp Ala Arg Arg Arg Arg Arg Gly
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Ala Ala Asp Asp His Ala Gly Val Val Gln Val Leu Ile Asp Phe Ala
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Glu Lys Arg Arg Gly His Ala His His Arg Lys Ala Gly Ala Met Asn						
		195		200		205
Ala Pro Phe Ile Leu Asp Leu Asp Cys Asp Tyr Tyr Val Asn Asn Ser						
		210		215		220
Gln Ala Leu Arg Ala Gly Ile Cys Phe Met Ile Glu Arg Gly Gly Gly						
		225		230		240
Gly Ala Ala Glu Asp Ala Gly Ala Val Ala Phe Val Gln Phe Pro Gln						
		245		250		255
Arg Val Asp Gly Val Asp Pro Gly Asp Arg Tyr Ala Asn His Asn Arg						
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Val Leu Phe Asp Cys Thr Glu Leu Gly Leu Asp Gly Leu Gln Gly Pro						
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Ile Tyr Val Gly Thr Gly Cys Leu Phe Arg Arg Val Ala Leu Tyr Ser						
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Val Asp Leu Pro Arg Trp Arg Pro Arg Arg Ser Leu Gly Cys Arg Leu						
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 <213> Glycine max

<220>
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 <222> (201)
 <223> Xaa = any amino acid

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Asp	Gly	Gln	Val	Cys	Glu	Ile	Cys	Gly	Asp	Gly	Val	Gly	Leu	Thr	Val
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Gln	Cys	Lys	Thr	Arg	Tyr	Lys	Arg	Leu	Lys	Gly	Ser	Pro	Arg	Val	Glu
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Gly	Asp	Asp	Asp	Glu	Glu	Asp	Val	Asp	Asp	Ile	Glu	His	Glu	Phe	Asn
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His	Gly	Arg	Met	Ser	Tyr	Gly	Arg	Gly	Pro	Glu	Asp	Asp	Asp	Asn	Ser
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Ser	Ser	Leu	His	Lys	Arg	Val	His	Pro	Tyr	Pro	Val	Ser	Glu	Pro	Gly
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Glu	Ile	Trp	Phe	Ala	Phe	Ser	Trp	Ile	Leu	Asp	Gln	Phe	Pro	Lys	Trp
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Ser	Ile	Glu	Pro	Arg	Ala	Pro	Glu	Met	Tyr	Phe	Ser	Glu	Lys	Ile	Asp	405	410	415	
Tyr	Leu	Lys	Asp	Lys	Val	Gln	Pro	Thr	Phe	Val	Lys	Glu	Arg	Arg	Ala	420	425	430	
Met	Lys	Arg	Glu	Tyr	Glu	Glu	Phe	Lys	Val	Arg	Ile	Asn	Ala	Leu	Val	435	440	445	
Ala	Lys	Ala	Gln	Lys	Val	Pro	Gln	Gly	Gly	Trp	Ile	Met	Gln	Asp	Gly	450	455	460	
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His	Lys	Lys	Ala	Gly	Ala	Met	Asn	Ala	Leu	Val	Arg	Val	Ser	Ala	Val	515	520	525	
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Asn	Asn	Ser	Lys	Ala	Ala	Arg	Glu	Ala	Met	Cys	Phe	Leu	Met	Asp	Pro	545	550	555	560
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Thr	Val	Tyr	Pro 820	Phe	Thr	Ser	Ile	Pro 825	Leu	Val	Ala	Tyr	Cys 830	Ile	Leu
Pro	Ala	Val 835	Cys	Leu	Leu	Thr	Asp 840	Lys	Phe	Ile	Met	Pro 845	Pro	Ile	Ser
Thr 850	Phe	Ala	Gly	Leu	Tyr	Phe 855	Val	Ala	Leu	Phe	Ser 860	Ser	Ile	Ile	Ala
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Phe	Ala	Val	Ile 900	Gln	Gly	Leu	Leu	Lys 905	Val	Leu	Ala	Gly	Ile 910	Asp	Thr

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 945 950 955 960
 Asn Asn Gly Tyr Gln Ser Trp Gly Pro Leu Phe Gly Lys Leu Phe Phe
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 Ser Phe Trp Val Ile Val His Leu Tyr Pro Phe Leu Lys Gly Leu Met
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 995 1000 1005
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 <212> DNA
 <213> Glycine max

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<210> 16
<211> 610
<212> PRT
<213> Glycine max

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<400> 16

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Val Phe Leu Gly His Ser Gly Gly Leu Asp Thr Asp Gly Asn Glu Leu
      35              40              45

Pro Arg Leu Val Tyr Val Ser Arg Glu Lys Arg Pro Gly Phe Gln His
      50              55              60

His Lys Lys Ala Gly Ala Met Asn Ala Leu Ile Arg Val Ser Ala Val
      65              70              75              80

Leu Thr Asn Gly Ala Tyr Leu Leu Asn Val Asp Cys Asp His Tyr Phe
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Asn Asn Ser Lys Ala Leu Lys Glu Ala Met Cys Phe Met Met Asp Pro
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Val Leu Gly Lys Lys Thr Cys Tyr Val Gln Phe Pro Gln Arg Phe Asp
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Gly Ile Asp Leu His Asp Arg Tyr Ala Asn Arg Asn Ile Val Phe Phe
      130             135             140

Asp Ile Asn Met Lys Gly Gln Asp Gly Val Gln Gly Pro Val Tyr Val
      145             150             155             160

Gly Thr Gly Cys Cys Phe Asn Arg Gln Ala Leu Tyr Gly Tyr Asp Pro
      165             170             175

Val Leu Thr Glu Glu Asp Leu Glu Pro Asn Ile Ile Val Lys Ser Cys
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Cys Gly Ser Arg Lys Lys Gly Lys Gly Gly Asn Lys Lys Tyr Ser Asp
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Lys Lys Lys Ala Met Gly Arg Thr Glu Ser Thr Val Pro Ile Phe Asn

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Val	Phe	Ile	Ala	Ala	Thr	Phe	Met	Glu	Gln	Gly	Gly	Ile	Pro	Pro	Ser
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Thr	Asn	Pro	Ala	Thr	Leu	Leu	Lys	Glu	Ala	Ile	His	Val	Ile	Ser	Cys
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Gly	Trp	Ile	Ser	Ile	Tyr	Cys	Met	Pro	Pro	Arg	Pro	Ala	Phe	Lys	Gly
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Tyr	Gly	Tyr	Asn	Gly	Lys	Leu	Lys	Pro	Leu	Met	Arg	Leu	Ala	Tyr	Ile
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Thr	Leu	Pro	Ala	Phe	Cys	Leu	Leu	Thr	Asn	Lys	Phe	Ile	Ile	Pro	Glu
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Ile	Ser	Asn	Phe	Ala	Ser	Met	Trp	Phe	Ile	Leu	Leu	Phe	Val	Ser	Ile
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Phe	Thr	Thr	Ser	Ile	Leu	Glu	Leu	Arg	Trp	Ser	Gly	Val	Ser	Ile	Glu
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His	Leu	Phe	Ala	Val	Phe	Gln	Gly	Leu	Leu	Lys	Val	Leu	Ala	Gly	Ile
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Asp	Thr	Asn	Phe	Thr	Val	Thr	Ser	Lys	Ala	Ser	Asp	Glu	Asp	Gly	Asp
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Phe	Ala	Glu	Leu	Tyr	Val	Phe	Lys	Trp	Thr	Ser	Leu	Leu	Ile	Pro	Pro
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Leu Phe Phe Ala Ile Trp Val Ile Ala His Leu Tyr Pro Phe Leu Lys				
545		550		555
Gly Leu Leu Gly Arg Gln Asn Arg Thr Pro Thr Ile Val Ile Val Trp				
		565		570
Ser Val Leu Leu Ala Ser Ile Phe Ser Leu Leu Trp Val Arg Ile Asp				
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Pro Phe Thr Ser Asp Ser Asn Lys Leu Thr Asn Gly Gln Cys Gly Ile				
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Asn Cys				
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 <211> 2890
 <212> DNA
 <213> Glycine max

<400> 17

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ggaacgattt	tcctacatta	actcgggtcgt	atatccctgg	acttccctcc	cattgcttgt	1740
ctactgtact	ctaccagcca	tatgccttct	gactggaaaa	tttatcgtac	ccgagattag	1800
caactatgcc	agtcttgtgt	tcattggccct	cttcatatcc	attgcagcaa	ctggcatcct	1860

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tgagatgcaa tggggcggtg ttagcataga cgactgggtg aggaacgaac agttttgggt 1920
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<210> 18
<211> 793
<212> PRT
<213> Glycine max

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Val Ile Cys Glu Ile Trp Phe Ala Val Ser Trp Ile Met Asp Gln Phe
      20              25              30

Pro Lys Trp Tyr Pro Ile Gln Arg Glu Thr Tyr Leu Asp Arg Leu Ser
      35              40              45

Leu Arg Tyr Glu Lys Glu Gly Lys Pro Ser Glu Leu Ser Ser Val Asp
      50              55              60

Val Phe Val Ser Thr Val Asp Pro Met Lys Glu Pro Pro Leu Ile Thr
      65              70              75              80

Ala Asn Thr Val Leu Ser Ile Leu Ala Val Asp Tyr Pro Val Asp Lys
      85              90              95

Val Ala Cys Tyr Val Ser Asp Asp Gly Ala Ala Met Leu Thr Phe Glu
      100             105             110

Ala Leu Ser Glu Thr Ser Glu Phe Ala Arg Arg Trp Val Pro Phe Cys
      115             120             125

Lys Lys Tyr Asn Ile Glu Pro Arg Ala Pro Glu Trp Tyr Phe Gly Gln
      130             135             140

Lys Met Asp Tyr Leu Lys Asn Lys Val His Pro Ala Phe Val Arg Glu
      145             150             155             160

Arg Arg Ala Met Lys Arg Asp Tyr Glu Glu Phe Lys Val Arg Ile Asn
      165             170             175

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Ser	Leu	Val	Ala	Thr	Ala	Gln	Lys	Val	Pro	Glu	Asp	Gly	Trp	Thr	Met	180	185	190	
Gln	Asp	Gly	Thr	Pro	Trp	Pro	Gly	Asn	Asn	Val	Arg	Asp	His	Pro	Gly	195	200	205	
Met	Ile	Gln	Val	Phe	Leu	Gly	Gln	Asp	Gly	Val	Arg	Asp	Val	Glu	Gly	210	215	220	
Asn	Glu	Leu	Pro	Arg	Leu	Val	Tyr	Val	Ser	Arg	Glu	Lys	Arg	Pro	Gly	225	230	235	240
Phe	Asp	His	His	Lys	Lys	Ala	Gly	Ala	Met	Asn	Ala	Leu	Val	Arg	Ala	245	250	255	
Ser	Ala	Ile	Ile	Thr	Asn	Ala	Pro	Tyr	Leu	Leu	Asn	Val	Asp	Cys	Asp	260	265	270	
His	Tyr	Ile	Asn	Asn	Ser	Lys	Ala	Leu	Arg	Glu	Ala	Met	Cys	Phe	Met	275	280	285	
Met	Asp	Pro	Gln	Leu	Gly	Lys	Lys	Val	Cys	Tyr	Val	Gln	Phe	Pro	Gln	290	295	300	
Arg	Phe	Asp	Gly	Ile	Asp	Arg	His	Asp	Arg	Tyr	Ser	Asn	Arg	Asn	Val	305	310	315	320
Val	Phe	Phe	Asp	Ile	Asn	Met	Lys	Gly	Leu	Asp	Gly	Ile	Gln	Gly	Pro	325	330	335	
Ile	Tyr	Val	Gly	Thr	Gly	Cys	Val	Phe	Arg	Arg	Tyr	Ala	Leu	Tyr	Gly	340	345	350	
Tyr	Asp	Ala	Pro	Ala	Lys	Lys	Lys	Pro	Pro	Ser	Lys	Thr	Cys	Asn	Cys	355	360	365	
Trp	Pro	Lys	Trp	Cys	Cys	Leu	Cys	Cys	Gly	Ser	Arg	Lys	Lys	Lys	Asn	370	375	380	
Ala	Asn	Ser	Lys	Lys	Glu	Lys	Lys	Arg	Lys	Val	Lys	His	Ser	Glu	Ala	385	390	395	400
Ser	Lys	Gln	Ile	His	Ala	Leu	Glu	Asn	Ile	Glu	Ala	Gly	Asn	Glu	Gly	405	410	415	
Thr	Asn	Asn	Glu	Lys	Thr	Ser	Asn	Leu	Thr	Gln	Thr	Lys	Leu	Glu	Lys	420	425	430	
Arg	Phe	Gly	Gln	Ser	Pro	Val	Phe	Val	Ala	Ser	Thr	Leu	Leu	Asp	Asp	435	440	445	
Gly	Gly	Val	Pro	His	Gly	Val	Ser	Pro	Ala	Ser	Leu	Leu	Lys	Glu	Ala	450	455	460	
Ile	Gln	Val	Ile	Ser	Cys	Gly	Tyr	Glu	Asp	Lys	Thr	Glu	Trp	Gly	Lys	465	470	475	480

Glu	Val	Gly	Trp	Ile	Tyr	Gly	Ser	Val	Thr	Glu	Asp	Ile	Leu	Thr	Gly	485	490	495
Phe	Lys	Met	His	Cys	His	Gly	Trp	Arg	Ser	Val	Tyr	Cys	Ile	Pro	Lys	500	505	510
Arg	Pro	Ala	Phe	Lys	Gly	Ser	Ala	Pro	Ile	Asn	Leu	Ser	Asp	Arg	Leu	515	520	525
His	Gln	Val	Leu	Arg	Trp	Ala	Leu	Gly	Ser	Val	Glu	Ile	Phe	Phe	Ser	530	535	540
Arg	His	Cys	Pro	Ile	Trp	Tyr	Gly	Tyr	Gly	Gly	Gly	Leu	Lys	Leu	Leu	545	550	555
Glu	Arg	Phe	Ser	Tyr	Ile	Asn	Ser	Val	Val	Tyr	Pro	Trp	Thr	Ser	Leu	565	570	575
Pro	Leu	Leu	Val	Tyr	Cys	Thr	Leu	Pro	Ala	Ile	Cys	Leu	Leu	Thr	Gly	580	585	590
Lys	Phe	Ile	Val	Pro	Glu	Ile	Ser	Asn	Tyr	Ala	Ser	Leu	Val	Phe	Met	595	600	605
Ala	Leu	Phe	Ile	Ser	Ile	Ala	Ala	Thr	Gly	Ile	Leu	Glu	Met	Gln	Trp	610	615	620
Gly	Gly	Val	Ser	Ile	Asp	Asp	Trp	Trp	Arg	Asn	Glu	Gln	Phe	Trp	Val	625	630	635
Ile	Gly	Gly	Val	Ser	Ser	His	Leu	Phe	Ala	Leu	Phe	Gln	Gly	Leu	Leu	645	650	655
Lys	Val	Leu	Ala	Gly	Val	Asn	Thr	Asn	Phe	Thr	Val	Thr	Ser	Lys	Ala	660	665	670
Ala	Asp	Asp	Gly	Glu	Phe	Ser	Glu	Leu	Tyr	Ile	Phe	Lys	Trp	Thr	Ser	675	680	685
Leu	Leu	Ile	Pro	Pro	Met	Thr	Leu	Leu	Ile	Met	Asn	Ile	Val	Gly	Val	690	695	700
Val	Val	Gly	Ile	Ser	Asp	Ala	Ile	Asn	Asn	Gly	Tyr	Asp	Ser	Trp	Gly	705	710	715
Pro	Leu	Phe	Gly	Arg	Leu	Phe	Phe	Ala	Leu	Trp	Val	Ile	Leu	His	Leu	725	730	735
Tyr	Pro	Phe	Leu	Lys	Gly	Leu	Leu	Gly	Lys	Gln	Asp	Arg	Met	Pro	Thr	740	745	750
Ile	Ile	Leu	Val	Trp	Ser	Ile	Leu	Leu	Ala	Ser	Ile	Leu	Thr	Leu	Met	755	760	765
Trp	Val	Arg	Ile	Asn	Pro	Phe	Val	Ser	Arg	Asp	Gly	Pro	Val	Leu	Glu	770	775	780

Ile Cys Gly Leu Asn Cys Asp Glu Ser
785 790

<210> 19
<211> 1733
<212> DNA
<213> Triticum aestivum

<220>
<221> unsure
<222> (262)
<223> n = a, c, g or t

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ccacaaagggt ttgatgggtat tgataggaat gatcgatatg caaacaggaa cactgtcttt 120
tttgatatta acttgagggg ccttgacggc attcaaggac cagtttatgt gggaactggg 180
tgtgttttca acagaacggc tatctatggg tatgagcccc caattaaggc gaagaagcca 240
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gacatagagg aggggtgttga aggtgctggg tttgatgatg agaaatcagt tctcatgtct 420
caaattgagct tagagaagag atttgggccag tcagcagcat ttggtgcctc cactctgatg 480
gaatatgggtg gtgttcctca gtcgtccact ccagaatctc ttttgaaaga agctatccat 540
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ggatctgtca cagaagatat tctaactgga ttcaagatgc acgcaagagg ctggcggttca 660
atctattgca tgcccaagcg cccagctttc aagggatctg ccccatcaa tctttcagat 720
cgtctgaatc aagtgtctgc gtgggctctt gggtctgttg aaattctttt cagccggcat 780
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gagaaagttg tcaaaattga gaaaacacat ttgtaaatag atgtaataga ctatctaccg 1680
ttttcatgag gttaagctct tcttttttgg aaaaaaaaaa aaaaaaaaaa aaa 1733

<210> 20
<211> 506
<212> PRT
<213> Triticum aestivum

<220>
<221> UNSURE
<222> (88)
<223> Xaa = any amino acid

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Tyr Val Gln Phe Pro Gln Arg Phe Asp Gly Ile Asp Arg Asn Asp Arg	20	25	30
Tyr Ala Asn Arg Asn Thr Val Phe Phe Asp Ile Asn Leu Arg Gly Leu	35	40	45
Asp Gly Ile Gln Gly Pro Val Tyr Val Gly Thr Gly Cys Val Phe Asn	50	55	60
Arg Thr Ala Ile Tyr Gly Tyr Glu Pro Pro Ile Lys Ala Lys Lys Pro	65	70	75
Gly Phe Leu Ala Ser Leu Cys Xaa Gly Lys Lys Lys Ala Ser Lys Ser	85	90	95
Lys Lys Arg Ser Ser Asp Lys Lys Lys Ser Asn Lys His Val Asp Ser	100	105	110
Ser Val Pro Val Phe Asn Leu Glu Asp Ile Glu Glu Gly Val Glu Gly	115	120	125
Ala Gly Phe Asp Asp Glu Lys Ser Val Leu Met Ser Gln Met Ser Leu	130	135	140
Glu Lys Arg Phe Gly Gln Ser Ala Ala Phe Val Ala Ser Thr Leu Met	145	150	155
Glu Tyr Gly Gly Val Pro Gln Ser Ser Thr Pro Glu Ser Leu Leu Lys	165	170	175
Glu Ala Ile His Val Ile Ser Cys Gly Tyr Glu Asp Lys Ser Glu Trp	180	185	190
Gly Thr Glu Ile Gly Trp Ile Tyr Gly Ser Val Thr Glu Asp Ile Leu	195	200	205
Thr Gly Phe Lys Met His Ala Arg Gly Trp Arg Ser Ile Tyr Cys Met	210	215	220
Pro Lys Arg Pro Ala Phe Lys Gly Ser Ala Pro Ile Asn Leu Ser Asp	225	230	235
Arg Leu Asn Gln Val Leu Arg Trp Ala Leu Gly Ser Val Glu Ile Leu	245	250	255
Phe Ser Arg His Cys Pro Leu Trp Tyr Gly Tyr Gly Gly Arg Leu Lys	260	265	270
Phe Leu Glu Arg Phe Ala Tyr Ile Asn Thr Thr Ile Tyr Pro Leu Thr	275	280	285
Ser Leu Pro Leu Leu Val Tyr Cys Ile Leu Pro Ala Ile Cys Leu Leu	290	295	300
Thr Gly Lys Phe Ile Met Pro Glu Ile Ser Asn Leu Ala Ser Ile Trp			

305		310		315		320
Phe	Ile	Ala	Leu	Phe	Leu	Met
		325		330		335
Arg	Trp	Ser	Gly	Val	Gly	Ile
		340		345		350
Trp	Val	Ile	Gly	Gly	Ile	Ser
		355		360		365
Leu	Leu	Lys	Val	Leu	Ala	Gly
		370		375		380
Lys	Ala	Asn	Asp	Glu	Glu	Gly
385				390		395
Trp	Thr	Thr	Leu	Leu	Ile	Pro
			405			410
Val	Gly	Val	Val	Ala	Gly	Thr
		420			425	
Ser	Trp	Gly	Pro	Leu	Phe	Gly
		435			440	
Val	His	Leu	Tyr	Pro	Phe	Leu
		450			455	
Thr	Pro	Thr	Ile	Val	Ile	Val
465				470		475
Ser	Leu	Leu	Trp	Val	Arg	Val
			485			490
Pro	Asn	Ile	Gln	Thr	Cys	Gly
		500			505	

<210> 21
 <211> 1029
 <212> DNA
 <213> Triticum aestivum

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 ttctgcagat cccaaccttc cagtgcacc gagatccatg gaccggtcca aggatctggc 180
 cgcctacgga tatggcagcg tggcctggaa ggagagaatg gagggctgga agcagaagca 240
 ggagcgccctg cagcatgtca ggagcgaggg tggcggtgat tgggatggcg acgatgcaga 300
 tctgccacta atggatgaag ctaggcagcc attgtccaga aaagtcccta tatcatcaag 360
 ccgaattaat ccctacagga tgattatcgt tatccggttg gtggtttttg gtttcttctt 420
 ccactaccga gtgatgcac cggcgaaaga tgcatttgca ttgtggctca tatctgtaat 480
 ctgtgaaatc tggtttgca tgtcctgtat tcttgatcag ttcccaaagt ggtttccaat 540
 cgagagagag acttacctgg accgtttgtc actaagggtt gacaaggaag gtcaaccctc 600
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 ggtcacagcg aacactgtcc tttccatcct ttctgtggat tatccggttg agaaggcttc 720
 ctgctatggt tctgatgatg gtgctgcaat gcttacgttt gaagcattgt ctgaaacatc 780

tgaatttgca aagaaatggg ttcctttcag caaaaagttt aatatcgagc ctcgtgctcc 840
 tgagtggtag ttccaacaga agatagacta cctgaaagac aagggttgctg cttcatttgt 900
 tagggagagg agggcgatga agagagaata cgaggaattc aaggtaagga tcaatgcctt 960
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 gcctggaaa 1029

<210> 22
 <211> 340
 <212> PRT
 <213> Triticum aestivum

<400> 22
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 20 25 30
 Arg Ile His Pro Leu Pro Phe Ala Asp Pro Asn Leu Pro Val Gln Pro
 35 40 45
 Arg Ser Met Asp Pro Ser Lys Asp Leu Ala Ala Tyr Gly Tyr Gly Ser
 50 55 60
 Val Ala Trp Lys Glu Arg Met Glu Gly Trp Lys Gln Lys Gln Glu Arg
 65 70 75 80
 Leu Gln His Val Arg Ser Glu Gly Gly Gly Asp Trp Asp Gly Asp Asp
 85 90 95
 Ala Asp Leu Pro Leu Met Asp Glu Ala Arg Gln Pro Leu Ser Arg Lys
 100 105 110
 Val Pro Ile Ser Ser Ser Arg Ile Asn Pro Tyr Arg Met Ile Ile Val
 115 120 125
 Ile Arg Leu Val Val Leu Gly Phe Phe Phe His Tyr Arg Val Met His
 130 135 140
 Pro Ala Lys Asp Ala Phe Ala Leu Trp Leu Ile Ser Val Ile Cys Glu
 145 150 155 160
 Ile Trp Phe Ala Met Ser Cys Ile Leu Asp Gln Phe Pro Lys Trp Phe
 165 170 175
 Pro Ile Glu Arg Glu Thr Tyr Leu Asp Arg Leu Ser Leu Arg Phe Asp
 180 185 190
 Lys Glu Gly Gln Pro Ser Gln Leu Ala Pro Ile Asp Phe Phe Val Ser
 195 200 205
 Thr Val Asp Pro Thr Lys Glu Pro Pro Leu Val Thr Ala Asn Thr Val
 210 215 220
 Leu Ser Ile Leu Ser Val Asp Tyr Pro Val Glu Lys Val Ser Cys Tyr
 225 230 235 240

Val Ser Asp Asp Gly Ala Ala Met Leu Thr Phe Glu Ala Leu Ser Glu
 245 250 255
 Thr Ser Glu Phe Ala Lys Lys Trp Val Pro Phe Ser Lys Lys Phe Asn
 260 265 270
 Ile Glu Pro Arg Ala Pro Glu Trp Tyr Phe Gln Gln Lys Ile Asp Tyr
 275 280 285
 Leu Lys Asp Lys Val Ala Ala Ser Phe Val Arg Glu Arg Arg Ala Met
 290 295 300
 Lys Arg Glu Tyr Glu Glu Phe Lys Val Arg Ile Asn Ala Leu Val Ala
 305 310 315 320
 Lys Ala Gln Lys Val Pro Glu Glu Gly Trp Thr Met Gln Asp Gly Ser
 325 330 335
 Pro Trp Pro Gly
 340

<210> 23
 <211> 1081
 <212> PRT
 <213> Arabidopsis thaliana

<400> 23
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 Leu Val Arg Ile Arg His Glu Ser Asp Gly Gly Thr Lys Pro Leu Lys
 20 25 30
 Asn Met Asn Gly Gln Ile Cys Gln Ile Cys Gly Asp Asp Val Gly Leu
 35 40 45
 Ala Glu Thr Gly Asp Val Phe Val Ala Cys Asn Glu Cys Ala Phe Pro
 50 55 60
 Val Cys Arg Pro Cys Tyr Glu Tyr Glu Arg Lys Asp Gly Thr Gln Cys
 65 70 75 80
 Cys Pro Gln Cys Lys Thr Arg Phe Arg Arg His Arg Gly Ser Pro Arg
 85 90 95
 Val Glu Gly Asp Glu Asp Glu Asp Asp Val Asp Asp Ile Glu Asn Glu
 100 105 110
 Phe Asn Tyr Ala Gln Gly Ala Asn Lys Ala Arg His Gln Arg His Gly
 115 120 125
 Glu Glu Phe Ser Ser Ser Ser Arg His Glu Ser Gln Pro Ile Pro Leu
 130 135 140
 Leu Thr His Gly His Thr Val Ser Gly Glu Ile Arg Thr Pro Asp Thr
 145 150 155 160

Gln	Ser	Val	Arg	Thr	Thr	Ser	Gly	Pro	Leu	Gly	Pro	Ser	Asp	Arg	Asn	
				165					170					175		
Ala	Ile	Ser	Ser	Pro	Tyr	Ile	Asp	Pro	Arg	Gln	Pro	Val	Pro	Val	Arg	
			180					185					190			
Ile	Val	Asp	Pro	Ser	Lys	Asp	Leu	Asn	Ser	Tyr	Gly	Leu	Gly	Asn	Val	
		195					200					205				
Asp	Trp	Lys	Glu	Arg	Val	Glu	Gly	Trp	Lys	Leu	Lys	Gln	Glu	Lys	Asn	
	210					215					220					
Met	Leu	Gln	Met	Thr	Gly	Lys	Tyr	His	Glu	Gly	Lys	Gly	Gly	Glu	Ile	
225					230					235					240	
Glu	Gly	Thr	Gly	Ser	Asn	Gly	Glu	Glu	Leu	Gln	Met	Ala	Asp	Asp	Thr	
				245					250					255		
Arg	Leu	Pro	Met	Ser	Arg	Val	Val	Pro	Ile	Pro	Ser	Ser	Arg	Leu	Thr	
			260					265					270			
Pro	Tyr	Arg	Val	Val	Ile	Ile	Leu	Arg	Leu	Ile	Ile	Leu	Cys	Phe	Phe	
		275					280					285				
Leu	Gln	Tyr	Arg	Thr	Thr	His	Pro	Val	Lys	Asn	Ala	Tyr	Pro	Leu	Trp	
	290					295					300					
Leu	Thr	Ser	Val	Ile	Cys	Glu	Ile	Trp	Phe	Ala	Phe	Ser	Trp	Leu	Leu	
305					310					315					320	
Asp	Gln	Phe	Pro	Lys	Trp	Tyr	Pro	Ile	Asn	Arg	Glu	Thr	Tyr	Leu	Asp	
				325					330					335		
Arg	Leu	Ala	Ile	Arg	Tyr	Asp	Arg	Asp	Gly	Glu	Pro	Ser	Gln	Leu	Val	
			340					345					350			
Pro	Val	Asp	Val	Phe	Val	Ser	Thr	Val	Asp	Pro	Leu	Lys	Glu	Pro	Pro	
		355					360					365				
Leu	Val	Thr	Ala	Asn	Thr	Val	Leu	Ser	Ile	Leu	Ser	Val	Asp	Tyr	Pro	
	370					375					380					
Val	Asp	Lys	Val	Ala	Cys	Tyr	Val	Ser	Asp	Asp	Gly	Ser	Ala	Met	Leu	
385					390					395					400	
Thr	Phe	Glu	Ser	Leu	Ser	Glu	Thr	Ala	Glu	Phe	Ala	Lys	Lys	Trp	Val	
				405					410					415		
Pro	Phe	Cys	Lys	Lys	Phe	Asn	Ile	Glu	Pro	Arg	Ala	Pro	Glu	Phe	Tyr	
			420					425					430			
Phe	Ala	Gln	Lys	Ile	Asp	Tyr	Leu	Lys	Asp	Lys	Ile	Gln	Pro	Ser	Phe	
		435					440					445				
Val	Lys	Glu	Arg	Arg	Ala	Met	Lys	Arg	Glu	Tyr	Glu	Glu	Phe	Lys	Val	
	450					455					460					

Arg Ile Asn Ala Leu Val Ala Lys Ala Gln Lys Ile Pro Glu Glu Gly
 465 470 475 480
 Trp Thr Met Gln Asp Gly Thr Pro Trp Pro Gly Asn Asn Thr Arg Asp
 485 490 495
 His Pro Gly Met Ile Gln Val Phe Leu Gly His Ser Gly Gly Leu Asp
 500 505 510
 Thr Asp Gly Asn Glu Leu Pro Arg Leu Ile Tyr Val Ser Arg Glu Lys
 515 520 525
 Arg Pro Gly Phe Gln His His Lys Lys Ala Gly Ala Met Asn Ala Leu
 530 535 540
 Ile Arg Val Ser Ala Val Leu Thr Asn Gly Ala Tyr Leu Leu Asn Val
 545 550 555 560
 Asp Cys Asp His Tyr Phe Asn Asn Ser Lys Ala Ile Lys Glu Ala Met
 565 570 575
 Cys Phe Met Met Asp Pro Ala Ile Gly Lys Lys Cys Cys Tyr Val Gln
 580 585 590
 Phe Pro Gln Arg Phe Asp Gly Ile Asp Leu His Asp Arg Tyr Ala Asn
 595 600 605
 Arg Asn Ile Val Phe Phe Asp Ile Asn Met Lys Gly Leu Asp Gly Ile
 610 615 620
 Gln Gly Pro Val Tyr Val Gly Thr Gly Cys Cys Phe Asn Arg Gln Ala
 625 630 635 640
 Leu Tyr Gly Tyr Asp Pro Val Leu Thr Glu Glu Asp Leu Glu Pro Asn
 645 650 655
 Ile Ile Val Lys Ser Cys Cys Gly Ser Arg Lys Lys Gly Lys Ser Ser
 660 665 670
 Lys Lys Tyr Asn Tyr Glu Lys Arg Arg Gly Ile Asn Arg Ser Asp Ser
 675 680 685
 Asn Ala Pro Leu Phe Asn Met Glu Asp Ile Asp Glu Gly Phe Glu Gly
 690 695 700
 Tyr Asp Asp Glu Arg Ser Ile Leu Met Ser Gln Arg Ser Val Glu Lys
 705 710 715 720
 Arg Phe Gly Gln Ser Pro Val Phe Ile Ala Ala Thr Phe Met Glu Gln
 725 730 735
 Gly Gly Ile Pro Pro Thr Thr Asn Pro Ala Thr Leu Leu Lys Glu Ala
 740 745 750
 Ile His Val Ile Ser Cys Gly Tyr Glu Asp Lys Thr Glu Trp Gly Lys
 755 760 765

Glu Ile Gly Trp Ile Tyr Gly Ser Val Thr Glu Asp Ile Leu Thr Gly
 770 775 780
 Phe Lys Met His Ala Arg Gly Trp Ile Ser Ile Tyr Cys Asn Pro Pro
 785 790 795 800
 Arg Pro Ala Phe Lys Gly Ser Ala Pro Ile Asn Leu Ser Asp Arg Leu
 805 810 815
 Asn Gln Val Leu Arg Trp Ala Leu Gly Ser Ile Glu Ile Leu Leu Ser
 820 825 830
 Arg His Cys Pro Ile Trp Tyr Gly Tyr His Gly Arg Leu Arg Leu Leu
 835 840 845
 Glu Arg Ile Ala Tyr Ile Asn Thr Ile Val Tyr Pro Ile Thr Ser Ile
 850 855 860
 Pro Leu Ile Ala Tyr Cys Ile Leu Pro Ala Phe Cys Leu Ile Thr Asp
 865 870 875 880
 Arg Phe Ile Ile Pro Glu Ile Ser Asn Tyr Ala Ser Ile Trp Phe Ile
 885 890 895
 Leu Leu Phe Ile Ser Ile Ala Val Thr Gly Ile Leu Glu Leu Arg Trp
 900 905 910
 Ser Gly Val Ser Ile Glu Asp Trp Trp Arg Asn Glu Gln Phe Trp Val
 915 920 925
 Ile Gly Gly Thr Ser Ala His Leu Phe Ala Val Phe Gln Gly Leu Leu
 930 935 940
 Lys Val Leu Ala Gly Ile Asp Thr Asn Phe Thr Val Thr Ser Lys Ala
 945 950 955 960
 Thr Asp Glu Asp Gly Asp Phe Ala Glu Leu Tyr Ile Phe Lys Trp Thr
 965 970 975
 Ala Leu Leu Ile Pro Pro Thr Thr Val Leu Leu Val Asn Leu Ile Gly
 980 985 990
 Ile Val Ala Gly Val Ser Tyr Ala Val Asn Ser Gly Tyr Gln Ser Trp
 995 1000 1005
 Gly Pro Leu Phe Gly Lys Leu Phe Phe Ala Leu Trp Val Ile Ala His
 1010 1015 1020
 Leu Tyr Pro Phe Leu Lys Gly Leu Leu Gly Arg Gln Asn Arg Thr Pro
 1025 1030 1035 1040
 Thr Ile Val Ile Val Trp Ser Val Leu Leu Ala Ser Ile Phe Ser Leu
 1045 1050 1055
 Leu Trp Val Arg Ile Asn Pro Phe Val Asp Ala Asn Pro Asn Ala Asn
 1060 1065 1070

Asn Phe Asn Gly Lys Gly Gly Val Phe
 1075 1080

<210> 24

<211> 1084

<212> PRT

<213> Arabidopsis thaliana

<400> 24

Met Asn Thr Gly Gly Arg Leu Ile Ala Gly Ser His Asn Arg Asn Glu
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Phe Val Leu Ile Asn Ala Asp Glu Ser Ala Arg Ile Arg Ser Val Gln
 20 25 30

Glu Leu Ser Gly Gln Thr Cys Gln Ile Cys Gly Asp Glu Ile Glu Leu
 35 40 45

Thr Val Ser Ser Glu Leu Phe Val Ala Cys Asn Glu Cys Ala Phe Pro
 50 55 60

Val Cys Arg Pro Cys Tyr Glu Tyr Glu Arg Arg Glu Gly Asn Gln Ala
 65 70 75 80

Cys Pro Gln Cys Lys Thr Arg Tyr Lys Arg Ile Lys Gly Ser Pro Arg
 85 90 95

Val Asp Gly Asp Asp Glu Glu Glu Glu Asp Ile Asp Asp Leu Glu Tyr
 100 105 110

Glu Phe Asp His Gly Met Asp Pro Glu His Ala Ala Glu Ala Ala Leu
 115 120 125

Ser Ser Arg Leu Asn Thr Gly Arg Gly Gly Leu Asp Ser Ala Pro Pro
 130 135 140

Gly Ser Gln Ile Pro Leu Leu Thr Tyr Cys Asp Glu Asp Ala Asp Met
 145 150 155 160

Tyr Ser Asp Arg His Ala Leu Ile Val Pro Pro Ser Thr Gly Tyr Gly
 165 170 175

Asn Arg Val Tyr Pro Ala Pro Phe Thr Asp Ser Ser Ala Pro Pro Gln
 180 185 190

Ala Arg Ser Met Val Pro Gln Lys Asp Ile Ala Glu Tyr Gly Tyr Gly
 195 200 205

Ser Val Ala Trp Lys Asp Arg Met Glu Val Trp Lys Arg Arg Gln Gly
 210 215 220

Glu Lys Leu Gln Val Ile Lys His Glu Gly Gly Asn Asn Gly Arg Gly
 225 230 235 240

Ser Asn Asp Asp Asp Glu Leu Asp Asp Pro Asp Met Pro Met Met Asp
 245 250 255

Glu	Gly	Arg	Gln	Pro	Leu	Ser	Arg	Lys	Leu	Pro	Ile	Arg	Ser	Ser	Arg			
			260					265					270					
Ile	Asn	Pro	Tyr	Arg	Met	Leu	Ile	Leu	Cys	Arg	Leu	Ala	Ile	Leu	Gly			
		275					280					285						
Leu	Phe	Phe	His	Tyr	Arg	Ile	Leu	His	Pro	Val	Asn	Asp	Ala	Tyr	Gly			
	290					295					300							
Leu	Trp	Leu	Thr	Ser	Val	Ile	Cys	Glu	Ile	Trp	Phe	Ala	Val	Ser	Trp			
305					310					315					320			
Ile	Leu	Asp	Gln	Phe	Pro	Lys	Trp	Tyr	Pro	Ile	Glu	Arg	Glu	Thr	Tyr			
				325				330						335				
Leu	Asp	Arg	Leu	Ser	Leu	Arg	Tyr	Glu	Lys	Glu	Gly	Lys	Pro	Ser	Gly			
			340					345					350					
Leu	Ala	Pro	Val	Asp	Val	Phe	Val	Ser	Thr	Val	Asp	Pro	Leu	Lys	Glu			
		355					360					365						
Pro	Pro	Leu	Ile	Thr	Ala	Asn	Thr	Val	Leu	Ser	Ile	Leu	Ala	Val	Asp			
	370					375					380							
Tyr	Pro	Val	Asp	Lys	Val	Ala	Cys	Tyr	Val	Ser	Asp	Asp	Gly	Ala	Ala			
385					390					395					400			
Met	Leu	Thr	Phe	Glu	Ala	Leu	Ser	Asp	Thr	Ala	Glu	Phe	Ala	Arg	Lys			
				405					410					415				
Trp	Val	Pro	Phe	Cys	Lys	Lys	Phe	Asn	Ile	Glu	Pro	Arg	Ala	Pro	Glu			
			420					425					430					
Trp	Tyr	Phe	Ser	Gln	Lys	Met	Asp	Tyr	Leu	Lys	Asn	Lys	Val	His	Pro			
		435					440					445						
Ala	Phe	Val	Arg	Glu	Arg	Arg	Ala	Met	Lys	Arg	Asp	Tyr	Glu	Glu	Phe			
		450				455					460							
Lys	Val	Lys	Ile	Asn	Ala	Leu	Val	Ala	Thr	Ala	Gln	Lys	Val	Pro	Glu			
465					470				475						480			
Glu	Gly	Trp	Thr	Met	Gln	Asp	Gly	Thr	Pro	Trp	Pro	Gly	Asn	Asn	Val			
				485					490					495				
Arg	Asp	His	Pro	Gly	Met	Ile	Gln	Val	Phe	Leu	Gly	His	Ser	Gly	Val			
			500					505					510					
Arg	Asp	Thr	Asp	Gly	Asn	Glu	Leu	Pro	Arg	Leu	Val	Tyr	Val	Ser	Arg			
		515					520					525						
Glu	Lys	Arg	Pro	Gly	Phe	Asp	His	His	Lys	Lys	Ala	Gly	Ala	Met	Asn			
		530				535					540							
Ser	Leu	Ile	Arg	Val	Ser	Ala	Val	Leu	Ser	Asn	Ala	Pro	Tyr	Leu	Leu			
545					550					555					560			

Asn	Val	Asp	Cys	Asp	His	Tyr	Ile	Asn	Asn	Ser	Lys	Ala	Ile	Arg	Glu	565	570	575
Ser	Met	Cys	Phe	Met	Met	Asp	Pro	Gln	Ser	Gly	Lys	Lys	Val	Cys	Tyr	580	585	590
Val	Gln	Phe	Pro	Gln	Arg	Phe	Asp	Gly	Ile	Asp	Arg	His	Asp	Arg	Tyr	595	600	605
Ser	Asn	Arg	Asn	Val	Val	Phe	Phe	Asp	Ile	Asn	Met	Lys	Gly	Leu	Asp	610	615	620
Gly	Ile	Gln	Gly	Pro	Ile	Tyr	Val	Gly	Thr	Gly	Cys	Val	Phe	Arg	Arg	625	630	635
Gln	Ala	Leu	Tyr	Gly	Phe	Asp	Ala	Pro	Lys	Lys	Lys	Lys	Pro	Pro	Gly	645	650	655
Lys	Thr	Cys	Asn	Cys	Trp	Pro	Lys	Trp	Cys	Cys	Leu	Cys	Cys	Gly	Leu	660	665	670
Arg	Lys	Lys	Ser	Lys	Thr	Lys	Ala	Lys	Asp	Lys	Lys	Thr	Asn	Thr	Lys	675	680	685
Glu	Thr	Ser	Lys	Gln	Ile	His	Ala	Leu	Glu	Asn	Val	Asp	Glu	Gly	Val	690	695	700
Ile	Val	Pro	Val	Ser	Asn	Val	Glu	Lys	Arg	Ser	Glu	Ala	Thr	Gln	Leu	705	710	715
Lys	Leu	Glu	Lys	Lys	Phe	Gly	Gln	Ser	Pro	Val	Phe	Val	Ala	Ser	Ala	725	730	735
Val	Leu	Gln	Asn	Gly	Gly	Val	Pro	Arg	Asn	Ala	Ser	Pro	Ala	Cys	Leu	740	745	750
Leu	Arg	Glu	Ala	Ile	Gln	Val	Ile	Ser	Cys	Gly	Tyr	Glu	Asp	Lys	Thr	755	760	765
Glu	Trp	Gly	Lys	Glu	Ile	Gly	Trp	Ile	Tyr	Gly	Ser	Val	Thr	Glu	Asp	770	775	780
Ile	Leu	Thr	Gly	Phe	Lys	Met	His	Cys	His	Gly	Trp	Arg	Ser	Val	Tyr	785	790	795
Cys	Met	Pro	Lys	Arg	Ala	Ala	Phe	Lys	Gly	Ser	Ala	Pro	Ile	Asn	Leu	805	810	815
Ser	Asp	Arg	Leu	His	Gln	Val	Leu	Arg	Trp	Ala	Leu	Gly	Ser	Val	Glu	820	825	830
Ile	Phe	Leu	Ser	Arg	His	Cys	Pro	Ile	Trp	Tyr	Gly	Tyr	Gly	Gly	Gly	835	840	845
Leu	Lys	Trp	Leu	Glu	Arg	Phe	Ser	Tyr	Ile	Asn	Ser	Val	Val	Tyr	Pro	850	855	860

Trp Thr Ser Leu Pro Leu Ile Val Tyr Cys Ser Leu Pro Ala Val Cys
 865 870 875 880
 Leu Leu Thr Gly Lys Phe Ile Val Pro Glu Ile Ser Asn Tyr Ala Gly
 885 890 895
 Ile Leu Phe Met Leu Met Phe Ile Ser Ile Ala Val Thr Gly Ile Leu
 900 905 910
 Glu Met Gln Trp Gly Gly Val Gly Ile Asp Asp Trp Trp Arg Asn Glu
 915 920 925
 Gln Phe Trp Val Ile Gly Gly Ala Ser Ser His Leu Phe Ala Leu Phe
 930 935 940
 Gln Gly Leu Leu Lys Val Leu Ala Gly Val Asn Thr Asn Phe Thr Val
 945 950 955 960
 Thr Ser Lys Ala Ala Asp Asp Gly Ala Phe Ser Glu Leu Tyr Ile Phe
 965 970 975
 Lys Trp Thr Thr Leu Leu Ile Pro Pro Thr Thr Leu Leu Ile Ile Asn
 980 985 990
 Ile Ile Gly Val Ile Val Gly Val Ser Asp Ala Ile Ser Asn Gly Tyr
 995 1000 1005
 Asp Ser Trp Gly Pro Leu Phe Gly Arg Leu Phe Phe Ala Leu Trp Val
 1010 1015 1020
 Ile Val His Leu Tyr Pro Phe Leu Lys Gly Met Leu Gly Lys Gln Asp
 1025 1030 1035 1040
 Lys Met Pro Thr Ile Ile Val Val Trp Ser Ile Leu Leu Ala Ser Ile
 1045 1050 1055
 Leu Thr Leu Leu Trp Val Arg Val Asn Pro Phe Val Ala Lys Gly Gly
 1060 1065 1070
 Pro Val Leu Glu Ile Cys Gly Leu Asn Cys Gly Asn
 1075 1080
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 <211> 685
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 Pro Glu Phe Tyr Phe Asn Glu Lys Ile Asp Tyr Leu Lys Asp Lys Val
 20 25 30
 His Pro Ser Phe Val Lys Glu Arg Arg Ala Met Lys Arg Glu Tyr Glu
 35 40 45

Glu	Phe	Lys	Val	Arg	Ile	Asn	Ala	Leu	Val	Ala	Lys	Ala	Gln	Lys	Lys	
50						55					60					
Pro	Glu	Glu	Gly	Trp	Val	Met	Gln	Asp	Gly	Thr	Pro	Trp	Pro	Gly	Asn	
65					70					75					80	
Asn	Thr	Arg	Asp	His	Pro	Gly	Met	Ile	Gln	Val	Tyr	Leu	Gly	Ser	Ala	
				85					90					95		
Gly	Ala	Leu	Asp	Val	Asp	Gly	Lys	Glu	Leu	Pro	Arg	Leu	Val	Tyr	Val	
			100					105					110			
Ser	Arg	Glu	Lys	Arg	Pro	Gly	Tyr	Gln	His	His	Lys	Lys	Ala	Gly	Ala	
		115					120					125				
Glu	Asn	Ala	Leu	Val	Arg	Val	Ser	Ala	Val	Leu	Thr	Asn	Ala	Pro	Phe	
		130				135					140					
Ile	Leu	Asn	Leu	Asp	Cys	Asp	His	Tyr	Ile	Asn	Asn	Ser	Lys	Ala	Met	
145					150					155					160	
Arg	Glu	Ala	Met	Cys	Phe	Leu	Met	Asp	Pro	Gln	Phe	Gly	Lys	Lys	Leu	
				165					170						175	
Cys	Tyr	Val	Gln	Phe	Pro	Gln	Arg	Phe	Asp	Gly	Ile	Asp	Arg	His	Asp	
			180					185					190			
Arg	Tyr	Ala	Asn	Arg	Asn	Val	Val	Phe	Phe	Asp	Ile	Asn	Met	Leu	Gly	
		195					200					205				
Leu	Asp	Gly	Leu	Gln	Gly	Pro	Val	Tyr	Val	Gly	Thr	Gly	Cys	Val	Phe	
	210					215					220					
Asn	Arg	Gln	Ala	Leu	Tyr	Gly	Tyr	Asp	Pro	Pro	Val	Ser	Glu	Lys	Arg	
225					230					235					240	
Pro	Lys	Met	Thr	Cys	Asp	Cys	Trp	Pro	Ser	Trp	Cys	Cys	Cys	Cys	Cys	
				245					250					255		
Gly	Gly	Ser	Arg	Lys	Lys	Ser	Lys	Lys	Lys	Gly	Glu	Lys	Lys	Gly	Leu	
			260					265						270		
Leu	Gly	Gly	Leu	Leu	Tyr	Gly	Lys	Lys	Lys	Lys	Met	Met	Gly	Lys	Asn	
		275					280					285				
Tyr	Val	Lys	Lys	Gly	Ser	Ala	Pro	Val	Phe	Asp	Leu	Glu	Glu	Ile	Glu	
	290					295					300					
Glu	Gly	Leu	Glu	Gly	Tyr	Glu	Glu	Leu	Glu	Lys	Ser	Thr	Leu	Met	Ser	
305					310					315					320	
Gln	Lys	Asn	Phe	Glu	Lys	Arg	Phe	Gly	Gln	Ser	Pro	Val	Phe	Ile	Ala	
				325					330					335		
Ser	Thr	Leu	Met	Glu	Asn	Gly	Gly	Leu	Pro	Glu	Gly	Thr	Asn	Ser	Thr	
			340					345					350			

Ser Ile Phe Ser Leu Val Trp Val Arg Ile Asp Pro Phe Leu Pro Lys
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Gln Thr Gly Pro Val Leu Lys Gln Cys Gly Val Glu Cys
675 680 685

<210> 26

<211> 1111

<212> PRT

<213> Arabidopsis thaliana

<400> 26

Met Ala Ser Thr Pro Pro Gln Thr Ser Lys Lys Val Arg Asn Asn Ser
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Gly Ser Gly Gln Thr Val Lys Phe Ala Arg Arg Thr Ser Ser Gly Arg
20 25 30

Tyr Val Ser Leu Ser Arg Asp Asn Ile Glu Leu Ser Gly Glu Leu Ser
35 40 45

Gly Asp Tyr Ser Asn Tyr Thr Val His Ile Pro Pro Thr Pro Asp Asn
50 55 60

Gln Pro Met Ala Thr Lys Ala Glu Glu Gln Tyr Val Ser Asn Ser Leu
65 70 75 80

Phe Thr Gly Gly Phe Asn Ser Val Thr Arg Ala His Leu Met Asp Lys
85 90 95

Val Ile Asp Ser Asp Val Thr His Pro Gln Met Ala Gly Ala Lys Gly
100 105 110

Ser Ser Cys Ala Met Pro Ala Cys Asp Gly Asn Val Met Lys Asp Glu
115 120 125

Arg Gly Lys Asp Val Met Pro Cys Glu Cys Arg Phe Lys Ile Cys Arg
130 135 140

Asp Cys Phe Met Asp Ala Gln Lys Glu Thr Gly Leu Cys Pro Gly Cys
145 150 155 160

Lys Glu Gln Tyr Lys Ile Gly Asp Leu Asp Asp Asp Thr Pro Asp Tyr
165 170 175

Ser Ser Gly Ala Leu Pro Leu Pro Ala Pro Gly Lys Asp Gln Arg Gly
180 185 190

Asn Asn Asn Asn Met Ser Met Met Lys Arg Asn Gln Asn Gly Glu Phe
195 200 205

Asp His Asn Arg Trp Leu Phe Glu Thr Gln Gly Thr Tyr Gly Tyr Gly
210 215 220

Asn Ala Tyr Trp Pro Gln Asp Glu Met Tyr Gly Asp Asp Met Asp Glu
225 230 235 240

Gly	Met	Arg	Gly	Gly	Met	Val	Glu	Thr	Ala	Asp	Lys	Pro	Trp	Arg	Pro	245	250	255
Leu	Ser	Arg	Arg	Ile	Pro	Ile	Pro	Ala	Ala	Ile	Ile	Ser	Pro	Tyr	Arg	260	265	270
Leu	Leu	Ile	Val	Ile	Arg	Phe	Val	Val	Leu	Cys	Phe	Phe	Leu	Thr	Trp	275	280	285
Arg	Ile	Arg	Asn	Pro	Asn	Glu	Asp	Ala	Ile	Trp	Leu	Trp	Leu	Met	Ser	290	295	300
Ile	Ile	Cys	Glu	Leu	Trp	Phe	Gly	Phe	Ser	Trp	Ile	Leu	Asp	Gln	Ile	305	310	315
Pro	Lys	Leu	Cys	Pro	Ile	Asn	Arg	Ser	Thr	Asp	Leu	Glu	Val	Leu	Arg	325	330	335
Asp	Lys	Phe	Asp	Met	Pro	Ser	Pro	Ser	Asn	Pro	Thr	Gly	Arg	Ser	Asp	340	345	350
Leu	Pro	Gly	Ile	Asp	Leu	Phe	Val	Ser	Thr	Ala	Asp	Pro	Glu	Lys	Glu	355	360	365
Pro	Pro	Leu	Val	Thr	Ala	Asn	Thr	Ile	Leu	Ser	Ile	Leu	Ala	Val	Asp	370	375	380
Tyr	Pro	Val	Glu	Lys	Val	Ser	Cys	Tyr	Leu	Ser	Asp	Asp	Gly	Gly	Ala	385	390	395
Leu	Leu	Ser	Phe	Glu	Ala	Met	Ala	Glu	Ala	Ala	Ser	Phe	Ala	Asp	Leu	405	410	415
Trp	Val	Pro	Phe	Cys	Arg	Lys	His	Asn	Ile	Glu	Pro	Arg	Asn	Pro	Asp	420	425	430
Ser	Tyr	Phe	Ser	Leu	Lys	Ile	Asp	Pro	Thr	Lys	Asn	Lys	Ser	Arg	Ile	435	440	445
Asp	Phe	Val	Lys	Asp	Arg	Arg	Lys	Ile	Lys	Arg	Glu	Tyr	Asp	Glu	Phe	450	455	460
Lys	Val	Arg	Ile	Asn	Gly	Leu	Pro	Asp	Ser	Ile	Arg	Arg	Arg	Ser	Asp	465	470	475
Ala	Phe	Asn	Ala	Arg	Glu	Glu	Met	Lys	Ala	Leu	Lys	Gln	Met	Arg	Glu	485	490	495
Ser	Gly	Gly	Asp	Pro	Thr	Glu	Pro	Val	Lys	Val	Pro	Lys	Ala	Thr	Trp	500	505	510
Met	Ala	Asp	Gly	Thr	His	Trp	Pro	Gly	Thr	Trp	Ala	Ala	Ser	Thr	Arg	515	520	525
Glu	His	Ser	Lys	Gly	Asp	His	Ala	Gly	Ile	Leu	Gln	Val	Met	Leu	Lys	530	535	540

Pro	Pro	Ser	Ser	Asp	Pro	Leu	Ile	Gly	Asn	Ser	Asp	Asp	Lys	Val	Ile	545	550	555	560
Asp	Phe	Ser	Asp	Thr	Asp	Thr	Arg	Leu	Pro	Met	Phe	Val	Tyr	Val	Ser		565	570	575
Arg	Glu	Lys	Arg	Pro	Gly	Tyr	Asp	His	Asn	Lys	Lys	Ala	Gly	Ala	Met	580	585	590	
Asn	Ala	Leu	Val	Arg	Ala	Ser	Ala	Ile	Leu	Ser	Asn	Gly	Pro	Phe	Ile	595	600	605	
Leu	Asn	Leu	Asp	Cys	Asp	His	Tyr	Ile	Tyr	Asn	Cys	Lys	Ala	Val	Arg	610	615	620	
Glu	Gly	Met	Cys	Phe	Met	Met	Asp	Arg	Gly	Gly	Glu	Asp	Ile	Cys	Tyr	625	630	635	640
Ile	Gln	Phe	Pro	Gln	Arg	Phe	Glu	Gly	Ile	Asp	Pro	Ser	Asp	Arg	Tyr		645	650	655
Ala	Asn	Asn	Asn	Thr	Val	Phe	Phe	Asp	Gly	Asn	Met	Arg	Ala	Leu	Asp	660	665	670	
Gly	Val	Gln	Gly	Pro	Val	Tyr	Val	Gly	Thr	Gly	Thr	Met	Phe	Arg	Arg		675	680	685
Phe	Ala	Leu	Tyr	Gly	Phe	Asp	Pro	Pro	Asn	Pro	Asp	Lys	Leu	Leu	Glu	690	695	700	
Lys	Lys	Glu	Ser	Glu	Thr	Glu	Ala	Leu	Thr	Thr	Ser	Asp	Phe	Asp	Pro	705	710	715	720
Asp	Leu	Asp	Val	Thr	Gln	Leu	Pro	Lys	Arg	Phe	Gly	Asn	Ser	Thr	Leu		725	730	735
Leu	Ala	Glu	Ser	Ile	Pro	Ile	Ala	Glu	Phe	Gln	Gly	Arg	Pro	Leu	Ala	740	745	750	
Asp	His	Pro	Ala	Val	Lys	Tyr	Gly	Arg	Pro	Pro	Gly	Ala	Leu	Arg	Val	755	760	765	
Pro	Arg	Asp	Pro	Leu	Asp	Ala	Thr	Thr	Val	Ala	Glu	Ser	Val	Ser	Val	770	775	780	
Ile	Ser	Cys	Trp	Tyr	Glu	Asp	Lys	Thr	Glu	Trp	Gly	Asp	Arg	Val	Gly	785	790	795	800
Trp	Ile	Tyr	Gly	Ser	Val	Thr	Glu	Asp	Val	Val	Thr	Gly	Tyr	Arg	Met		805	810	815
His	Asn	Arg	Gly	Trp	Arg	Ser	Val	Tyr	Cys	Ile	Thr	Lys	Arg	Asp	Ser	820	825	830	
Phe	Arg	Gly	Ser	Ala	Pro	Ile	Asn	Leu	Thr	Asp	Arg	Leu	His	Gln	Val	835	840	845	

Met	Glu	Ala	Ser	Ala	Gly	Leu	Val	Ala	Gly	Ser	His	Asn	Arg	Asn	Glu	1	5	10	15
Leu	Val	Val	Ile	His	Asn	His	Glu	Glu	Pro	Lys	Pro	Leu	Lys	Asn	Leu	20	25	30	
Asp	Gly	Gln	Phe	Cys	Glu	Ile	Cys	Gly	Asp	Gln	Ile	Gly	Leu	Thr	Val	35	40	45	
Glu	Gly	Asp	Leu	Phe	Val	Ala	Cys	Asn	Glu	Cys	Gly	Phe	Pro	Ala	Cys	50	55	60	
Arg	Pro	Cys	Tyr	Glu	Tyr	Glu	Arg	Arg	Glu	Gly	Thr	Gln	Asn	Cys	Pro	65	70	75	80
Gln	Cys	Lys	Thr	Arg	Tyr	Lys	Arg	Leu	Arg	Gly	Ser	Pro	Arg	Val	Glu	85	90	95	
Gly	Asp	Glu	Asp	Glu	Glu	Asp	Ile	Asp	Asp	Ile	Glu	Tyr	Glu	Phe	Asn	100	105	110	
Ile	Glu	His	Glu	Gln	Asp	Lys	His	Lys	His	Ser	Ala	Glu	Ala	Met	Leu	115	120	125	
Tyr	Gly	Lys	Met	Ser	Tyr	Gly	Arg	Gly	Pro	Glu	Asp	Asp	Glu	Asn	Gly	130	135	140	
Arg	Phe	Pro	Pro	Val	Ile	Ala	Gly	Gly	His	Ser	Gly	Glu	Phe	Pro	Val	145	150	155	160
Gly	Gly	Gly	Tyr	Gly	Asn	Gly	Glu	His	Gly	Leu	His	Lys	Arg	Val	His	165	170	175	
Pro	Tyr	Pro	Ser	Ser	Glu	Ala	Gly	Ser	Glu	Gly	Gly	Trp	Arg	Glu	Arg	180	185	190	
Met	Asp	Asp	Trp	Lys	Leu	Gln	His	Gly	Asn	Leu	Gly	Pro	Glu	Pro	Asp	195	200	205	
Asp	Asp	Pro	Glu	Met	Gly	Leu	Ile	Asp	Glu	Ala	Arg	Gln	Pro	Leu	Ser	210	215	220	
Arg	Lys	Val	Pro	Ile	Ala	Ser	Ser	Lys	Ile	Asn	Pro	Tyr	Arg	Met	Val	225	230	235	240
Ile	Val	Ala	Arg	Leu	Val	Ile	Leu	Ala	Val	Phe	Leu	Arg	Tyr	Arg	Leu	245	250	255	
Leu	Asn	Pro	Val	His	Asp	Ala	Leu	Gly	Leu	Trp	Leu	Thr	Ser	Val	Ile	260	265	270	
Cys	Glu	Ile	Trp	Phe	Ala	Val	Ser	Trp	Ile	Leu	Asp	Gln	Phe	Pro	Lys	275	280	285	
Trp	Phe	Pro	Ile	Glu	Arg	Glu	Thr	Tyr	Leu	Asp	Arg	Leu	Ser	Leu	Arg	290	295	300	

Tyr	Glu	Arg	Glu	Gly	Glu	Pro	Asn	Met	Leu	Ala	Pro	Val	Asp	Val	Phe	305	310	315	320
Val	Ser	Thr	Val	Asp	Pro	Leu	Lys	Glu	Pro	Pro	Leu	Val	Thr	Ser	Asn	325	330	335	
Thr	Val	Leu	Ser	Ile	Leu	Ala	Met	Asp	Tyr	Pro	Val	Glu	Lys	Ile	Ser	340	345	350	
Cys	Tyr	Val	Ser	Asp	Asp	Gly	Ala	Ser	Met	Leu	Thr	Phe	Glu	Ser	Leu	355	360	365	
Ser	Glu	Thr	Ala	Glu	Phe	Ala	Arg	Lys	Trp	Val	Pro	Phe	Cys	Lys	Lys	370	375	380	
Phe	Ser	Ile	Glu	Pro	Arg	Ala	Pro	Glu	Met	Tyr	Phe	Thr	Leu	Lys	Val	385	390	395	400
Asp	Tyr	Leu	Gln	Asp	Lys	Val	His	Pro	Thr	Phe	Val	Lys	Glu	Arg	Arg	405	410	415	
Ala	Met	Lys	Arg	Glu	Tyr	Glu	Glu	Phe	Lys	Val	Arg	Ile	Asn	Ala	Gln	420	425	430	
Val	Ala	Lys	Ala	Ser	Lys	Val	Pro	Leu	Glu	Gly	Trp	Ile	Met	Gln	Asp	435	440	445	
Gly	Thr	Pro	Trp	Pro	Gly	Asn	Asn	Thr	Lys	Asp	His	Pro	Gly	Met	Ile	450	455	460	
Gln	Val	Phe	Leu	Gly	His	Ser	Gly	Gly	Phe	Asp	Val	Glu	Gly	His	Glu	465	470	475	480
Leu	Pro	Arg	Leu	Val	Tyr	Val	Ser	Arg	Glu	Lys	Arg	Pro	Gly	Phe	Gln	485	490	495	
His	His	Lys	Lys	Ala	Gly	Ala	Met	Asn	Ala	Leu	Val	Arg	Val	Ala	Gly	500	505	510	
Val	Leu	Thr	Asn	Ala	Pro	Phe	Met	Leu	Asn	Leu	Asp	Cys	Asp	His	Tyr	515	520	525	
Val	Asn	Asn	Ser	Lys	Ala	Val	Arg	Glu	Ala	Met	Cys	Phe	Leu	Met	Asp	530	535	540	
Pro	Gln	Ile	Gly	Lys	Lys	Val	Cys	Tyr	Val	Gln	Phe	Pro	Gln	Arg	Phe	545	550	555	560
Asp	Gly	Ile	Asp	Thr	Asn	Asp	Arg	Tyr	Ala	Asn	Arg	Asn	Thr	Val	Phe	565	570	575	
Phe	Asp	Ile	Asn	Met	Lys	Gly	Leu	Asp	Gly	Ile	Gln	Gly	Pro	Val	Tyr	580	585	590	
Val	Gly	Thr	Gly	Cys	Val	Phe	Lys	Arg	Gln	Ala	Leu	Tyr	Gly	Tyr	Glu	595	600	605	

Pro	Pro	Lys	Gly	Pro	Lys	Arg	Pro	Lys	Met	Ile	Ser	Cys	Gly	Cys	Cys	
610						615					620					
Pro	Cys	Phe	Gly	Arg	Arg	Arg	Lys	Asn	Lys	Lys	Phe	Ser	Lys	Asn	Asp	
625					630					635					640	
Met	Asn	Gly	Asp	Val	Ala	Ala	Leu	Gly	Gly	Ala	Glu	Gly	Asp	Lys	Glu	
				645					650					655		
His	Leu	Met	Phe	Glu	Met	Asn	Phe	Glu	Lys	Thr	Phe	Gly	Gln	Ser	Ser	
			660					665					670			
Ile	Phe	Val	Thr	Ser	Thr	Leu	Met	Glu	Glu	Gly	Gly	Val	Pro	Pro	Ser	
		675					680					685				
Ser	Ser	Pro	Ala	Val	Leu	Leu	Lys	Glu	Ala	Ile	His	Val	Ile	Ser	Cys	
		690				695					700					
Gly	Tyr	Glu	Asp	Lys	Thr	Glu	Trp	Gly	Thr	Glu	Leu	Gly	Trp	Ile	Tyr	
705					710					715					720	
Gly	Ser	Ile	Thr	Glu	Asp	Ile	Leu	Thr	Gly	Phe	Lys	Met	His	Cys	Arg	
				725					730					735		
Gly	Trp	Arg	Ser	Ile	Tyr	Cys	Met	Pro	Lys	Arg	Pro	Ala	Phe	Lys	Gly	
			740					745					750			
Ser	Ala	Pro	Ile	Asn	Leu	Ser	Asp	Arg	Leu	Asn	Gln	Val	Leu	Arg	Trp	
		755					760					765				
Ala	Leu	Gly	Ser	Val	Glu	Ile	Phe	Phe	Ser	Arg	His	Ser	Pro	Leu	Trp	
		770				775					780					
Tyr	Gly	Tyr	Lys	Gly	Gly	Lys	Leu	Lys	Trp	Leu	Glu	Arg	Phe	Ala	Tyr	
785					790					795					800	
Ala	Asn	Thr	Thr	Ile	Tyr	Pro	Phe	Thr	Ser	Ile	Pro	Leu	Leu	Ala	Tyr	
				805					810					815		
Cys	Ile	Leu	Pro	Ala	Ile	Cys	Leu	Leu	Thr	Asp	Lys	Phe	Ile	Met	Pro	
			820					825					830			
Pro	Ile	Ser	Thr	Phe	Ala	Ser	Leu	Phe	Phe	Ile	Ser	Leu	Phe	Met	Ser	
		835					840					845				
Ile	Ile	Val	Thr	Gly	Ile	Leu	Glu	Leu	Arg	Trp	Ser	Gly	Val	Ser	Ile	
		850				855					860					
Glu	Glu	Trp	Trp	Arg	Asn	Glu	Gln	Phe	Trp	Val	Ile	Gly	Gly	Ile	Ser	
865					870					875					880	
Ala	His	Leu	Phe	Ala	Val	Val	Gln	Gly	Leu	Leu	Lys	Ile	Leu	Ala	Gly	
				885					890					895		
Ile	Asp	Thr	Asn	Phe	Thr	Val	Thr	Ser	Lys	Ala	Thr	Asp	Asp	Asp	Asp	
			900					905					910			

Phe Gly Glu Leu Tyr Ala Phe Lys Trp Thr Thr Leu Leu Ile Pro Pro
 915 920 925
 Thr Thr Val Leu Ile Ile Asn Ile Val Gly Val Val Ala Gly Ile Ser
 930 935 940
 Asp Ala Ile Asn Asn Gly Tyr Gln Ser Trp Gly Pro Leu Phe Gly Lys
 945 950 955 960
 Leu Phe Phe Ser Phe Trp Val Ile Val His Leu Tyr Pro Phe Leu Lys
 965 970 975
 Gly Leu Met Gly Arg Gln Asn Arg Thr Pro Thr Ile Val Val Ile Trp
 980 985 990
 Ser Val Leu Leu Ala Ser Ile Phe Ser Leu Leu Trp Val Arg Ile Asp
 995 1000 1005
 Pro Phe Val Leu Lys Thr Lys Gly Pro Asp Thr Ser Lys Cys Gly Ile
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 Asn Cys
 1025
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 Ala Met Leu Thr Phe Glu Ala Leu Ser Glu Thr Ser Glu Phe Ala Arg
 20 25 30
 Lys Trp Val Pro Phe Cys Lys Lys Tyr Asn Ile Glu Pro Arg Ala Pro
 35 40 45
 Glu Trp Tyr Phe Ala Gln Lys Ile Asp Tyr Leu Lys Asp Lys Val Gln
 50 55 60
 Thr Ser Phe Val Lys Glu Arg Arg Ala Met Lys Arg Glu Tyr Glu Glu
 65 70 75 80
 Phe Lys Val Arg Val Asn Gly Leu Val Ala Lys Ala Gln Lys Val Pro
 85 90 95
 Glu Glu Gly Trp Ile Met Gln Asp Gly Thr Pro Trp Pro Gly Asn Asn
 100 105 110
 Thr Arg Asp His Pro Gly Met Ile Gln Val Phe Leu Gly Gln Ser Gly
 115 120 125
 Gly Leu Asp Ala Glu Gly Asn Glu Leu Pro Arg Leu Val Tyr Val Ser
 130 135 140

Arg	Glu	Lys	Arg	Pro	Gly	Phe	Gln	His	His	Lys	Lys	Ala	Gly	Ala	Met	145	150	155	160
Asn	Ala	Leu	Val	Arg	Val	Ser	Ala	Val	Leu	Thr	Asn	Gly	Ala	Phe	Leu	165	170		175
Leu	Asn	Leu	Asp	Cys	Asp	His	Tyr	Ile	Asn	Asn	Ser	Lys	Ala	Leu	Arg	180	185		190
Glu	Ala	Met	Cys	Phe	Leu	Met	Asp	Pro	Asn	Leu	Gly	Lys	Gln	Val	Cys	195	200	205	
Tyr	Val	Gln	Phe	Pro	Gln	Arg	Phe	Asp	Gly	Ile	Asp	Arg	Asn	Asp	Arg	210	215	220	
Tyr	Ala	Asn	Arg	Asn	Thr	Val	Phe	Phe	Asp	Ile	Asn	Leu	Arg	Gly	Leu	225	230	235	240
Asp	Gly	Ile	Gln	Gly	Pro	Val	Tyr	Val	Gly	Thr	Gly	Cys	Val	Phe	Asn	245	250		255
Arg	Thr	Ala	Leu	Tyr	Gly	Tyr	Glu	Pro	Pro	Leu	Lys	Pro	Lys	His	Arg	260	265	270	
Lys	Thr	Gly	Ile	Leu	Ser	Ser	Leu	Cys	Gly	Gly	Ser	Arg	Lys	Lys	Ser	275	280	285	
Ser	Lys	Ser	Ser	Lys	Lys	Gly	Ser	Asp	Lys	Lys	Lys	Ser	Gly	Lys	His	290	295	300	
Val	Asp	Ser	Thr	Val	Pro	Val	Phe	Asn	Leu	Glu	Asp	Ile	Glu	Glu	Gly	305	310	315	320
Val	Glu	Gly	Ala	Gly	Phe	Asp	Asp	Glu	Lys	Ser	Leu	Leu	Met	Ser	Gln	325	330		335
Met	Ser	Leu	Glu	Lys	Arg	Phe	Gly	Gln	Ser	Ala	Val	Phe	Val	Ala	Ser	340	345		350
Thr	Leu	Met	Glu	Asn	Gly	Gly	Val	Pro	Gln	Ser	Ala	Thr	Pro	Glu	Thr	355	360	365	
Leu	Leu	Lys	Glu	Ala	Ile	His	Val	Ile	Ser	Cys	Gly	Tyr	Glu	Asp	Lys	370	375	380	
Thr	Asp	Trp	Gly	Ser	Glu	Ile	Gly	Trp	Ile	Tyr	Gly	Ser	Val	Thr	Glu	385	390	395	400
Asp	Ile	Leu	Thr	Gly	Phe	Lys	Met	His	Ala	Arg	Gly	Trp	Arg	Ser	Ile	405	410		415
Tyr	Cys	Met	Pro	Lys	Arg	Pro	Ala	Phe	Lys	Gly	Ser	Ala	Pro	Ile	Asn	420	425	430	
Leu	Ser	Asp	Arg	Leu	Asn	Gln	Val	Leu	Arg	Trp	Ala	Leu	Gly	Ser	Val	435	440	445	

Glu Ile Leu Phe Ser Arg His Cys Pro Ile Trp Tyr Gly Tyr Ser Gly
 450 455 460
 Arg Leu Lys Trp Leu Glu Arg Phe Ala Tyr Val Asn Thr Thr Ile Tyr
 465 470 475 480
 Pro Val Thr Ala Ile Pro Leu Leu Met Tyr Cys Thr Leu Pro Ala Val
 485 490 495
 Cys Leu Leu Thr Asn Lys Phe Ile Ile Pro Gln Ile Ser Asn Leu Ala
 500 505 510
 Ser Ile Trp Phe Ile Ser Leu Phe Leu Ser Ile Phe Ala Thr Gly Ile
 515 520 525
 Leu Lys Met Lys Trp Asn Gly Val Gly Ile Asp Gln Trp Trp Arg Asn
 530 535 540
 Glu Gln Phe Trp Val Ile Gly Gly Val Ser Ala His Leu Phe Ala Val
 545 550 555 560
 Phe Gln Gly Leu Leu Lys Val Leu Ala Gly Ile Asp Thr Asn Phe Thr
 565 570 575
 Val Thr Ser Lys Ala Ser Asp Glu Asp Gly Asp Phe Ala Glu Leu Tyr
 580 585 590
 Met Phe Lys Trp Thr Thr Leu Leu Ile Pro Pro Thr Thr Leu Leu Ile
 595 600 605
 Ile Asn Leu Val Gly Val Val Ala Gly Ile Ser Tyr Val Ile Asn Ser
 610 615 620
 Gly Tyr Gln Ser Trp Gly Pro Leu Phe Gly Lys Leu Phe Phe Ala Phe
 625 630 635 640
 Trp Val Ile Ile His Leu Tyr Pro Phe Leu Lys Gly Leu Met Gly Arg
 645 650 655
 Gln Asn Arg Thr Pro Thr Ile Val Val Val Trp Ser Ile Leu Leu Ala
 660 665 670
 Ser Ile Phe Ser Leu Leu Trp Val Arg Ile Asp Pro Phe Thr Thr Arg
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 Val Thr Gly Pro Asp Val Glu Gln Cys Gly Ile Asn Cys
 690 695 700

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 <211> 1081
 <212> PRT
 <213> Arabidopsis thaliana

<400> 29
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Ile	Asn	Ala	Asp	Glu	Asn	Ala	Arg	Ile	Arg	Ser	Val	Gln	Glu	Leu	Ser	20	25	30	
Gly	Gln	Thr	Cys	Gln	Ile	Cys	Arg	Asp	Glu	Ile	Glu	Leu	Thr	Val	Asp	35	40	45	
Gly	Glu	Pro	Phe	Val	Ala	Cys	Asn	Glu	Cys	Ala	Phe	Pro	Val	Cys	Arg	50	55	60	
Pro	Cys	Tyr	Glu	Tyr	Glu	Arg	Arg	Glu	Gly	Asn	Gln	Ala	Cys	Pro	Gln	65	70	75	80
Cys	Lys	Thr	Arg	Phe	Lys	Arg	Leu	Lys	Gly	Ser	Pro	Arg	Val	Glu	Gly	85	90	95	
Asp	Glu	Glu	Glu	Asp	Asp	Ile	Asp	Asp	Leu	Asp	Asn	Glu	Phe	Glu	Tyr	100	105	110	
Gly	Asn	Asn	Gly	Ile	Gly	Phe	Asp	Gln	Val	Ser	Glu	Gly	Met	Ser	Ile	115	120	125	
Ser	Arg	Arg	Asn	Ser	Gly	Phe	Pro	Gln	Ser	Asp	Leu	Asp	Ser	Ala	Pro	130	135	140	
Pro	Gly	Ser	Gln	Ile	Pro	Leu	Leu	Thr	Tyr	Gly	Asp	Glu	Asp	Val	Glu	145	150	155	160
Ile	Ser	Ser	Asp	Arg	His	Ala	Leu	Ile	Val	Pro	Pro	Ser	Leu	Gly	Gly	165	170	175	
His	Gly	Asn	Arg	Val	His	Pro	Val	Ser	Leu	Ser	Asp	Pro	Thr	Val	Ala	180	185	190	
Ala	His	Arg	Arg	Leu	Met	Val	Pro	Gln	Lys	Asp	Leu	Ala	Val	Tyr	Gly	195	200	205	
Tyr	Gly	Ser	Val	Ala	Trp	Lys	Asp	Arg	Met	Glu	Glu	Trp	Lys	Arg	Lys	210	215	220	
Gln	Asn	Glu	Lys	Leu	Gln	Val	Val	Arg	His	Glu	Gly	Asp	Pro	Asp	Phe	225	230	235	240
Glu	Asp	Gly	Asp	Asp	Ala	Asp	Phe	Pro	Met	Met	Asp	Glu	Gly	Arg	Gln	245	250	255	
Pro	Leu	Ser	Met	Lys	Ile	Pro	Ile	Lys	Ser	Ser	Lys	Ile	Asn	Pro	Tyr	260	265	270	
Arg	Met	Leu	Ile	Val	Leu	Arg	Leu	Val	Ile	Leu	Gly	Leu	Phe	Phe	His	275	280	285	
Tyr	Arg	Ile	Leu	His	Pro	Val	Lys	Asp	Ala	Tyr	Ala	Leu	Trp	Leu	Ile	290	295	300	
Ser	Val	Ile	Cys	Glu	Ile	Trp	Phe	Ala	Val	Ser	Trp	Val	Leu	Asp	Gln	305	310	315	320

Phe Pro Lys Trp Tyr Pro Ile Glu Arg Glu Thr Tyr Leu Asp Arg Leu
325 330 335
Ser Leu Arg Tyr Glu Lys Glu Gly Lys Pro Ser Gly Leu Ser Pro Val
340 345 350
Asp Val Phe Val Ser Thr Val Asp Pro Leu Lys Glu Pro Pro Leu Ile
355 360 365
Thr Ala Asn Thr Val Leu Ser Ile Leu Ala Val Asp Tyr Pro Val Asp
370 375 380
Lys Val Ala Cys Tyr Val Ser Asp Asp Gly Ala Ala Met Leu Thr Phe
385 390 395 400
Glu Ala Leu Ser Glu Thr Ala Glu Phe Ala Arg Lys Trp Val Pro Phe
405 410 415
Cys Lys Lys Tyr Cys Ile Glu Pro Arg Ala Pro Glu Trp Tyr Phe Cys
420 425 430
His Lys Met Asp Tyr Leu Lys Asn Lys Val His Pro Ala Phe Val Arg
435 440 445
Glu Arg Arg Ala Met Lys Arg Asp Tyr Glu Glu Phe Lys Val Lys Ile
450 455 460
Asn Ala Leu Val Ala Thr Ala Gln Lys Val Pro Glu Asp Gly Trp Thr
465 470 475 480
Met Gln Asp Gly Thr Pro Trp Pro Gly Asn Ser Val Arg Asp His Pro
485 490 495
Gly Met Ile Gln Val Phe Leu Gly Ser Asp Gly Val Arg Asp Val Glu
500 505 510
Asn Asn Glu Leu Pro Arg Leu Val Tyr Val Ser Arg Glu Lys Arg Pro
515 520 525
Gly Phe Asp His His Lys Lys Ala Gly Ala Met Asn Ser Leu Ile Arg
530 535 540
Val Ser Gly Val Leu Ser Asn Ala Pro Tyr Leu Leu Asn Val Asp Cys
545 550 555 560
Asp His Tyr Ile Asn Asn Ser Lys Ala Leu Arg Glu Ala Met Cys Phe
565 570 575
Met Met Asp Pro Gln Ser Gly Lys Lys Ile Cys Tyr Val Gln Phe Pro
580 585 590
Gln Arg Phe Asp Gly Ile Asp Arg His Asp Arg Tyr Ser Asn Arg Asn
595 600 605
Val Val Phe Phe Asp Ile Asn Met Lys Gly Leu Asp Gly Leu Gln Gly
610 615 620

Pro	Ile	Tyr	Val	Gly	Thr	Gly	Cys	Val	Phe	Arg	Arg	Gln	Ala	Leu	Tyr	
625					630					635					640	
Gly	Phe	Asp	Ala	Pro	Lys	Lys	Lys	Lys	Gly	Pro	Arg	Lys	Thr	Cys	Asn	
				645					650					655		
Cys	Trp	Pro	Lys	Trp	Cys	Leu	Leu	Cys	Phe	Gly	Ser	Arg	Lys	Asn	Arg	
			660					665					670			
Lys	Ala	Lys	Thr	Val	Ala	Ala	Asp	Lys	Lys	Lys	Lys	Asn	Arg	Glu	Ala	
		675					680					685				
Ser	Lys	Gln	Ile	His	Ala	Leu	Glu	Asn	Ile	Glu	Glu	Gly	Arg	Gly	His	
	690					695				700						
Lys	Val	Leu	Asn	Val	Glu	Gln	Ser	Thr	Glu	Ala	Met	Gln	Met	Lys	Leu	
705					710					715					720	
Gln	Lys	Lys	Tyr	Gly	Gln	Ser	Pro	Val	Phe	Val	Ala	Ser	Ala	Arg	Leu	
				725					730					735		
Glu	Asn	Gly	Gly	Met	Ala	Arg	Asn	Ala	Ser	Pro	Ala	Cys	Leu	Leu	Lys	
			740					745					750			
Glu	Ala	Ile	Gln	Val	Ile	Ser	Arg	Gly	Tyr	Glu	Asp	Lys	Thr	Glu	Trp	
		755					760					765				
Gly	Lys	Glu	Ile	Gly	Trp	Ile	Tyr	Gly	Ser	Val	Thr	Glu	Asp	Ile	Leu	
	770					775					780					
Thr	Gly	Ser	Lys	Met	His	Ser	His	Gly	Trp	Arg	His	Val	Tyr	Cys	Thr	
785					790					795					800	
Pro	Lys	Leu	Ala	Ala	Phe	Lys	Gly	Ser	Ala	Pro	Ile	Asn	Leu	Ser	Asp	
				805					810					815		
Arg	Leu	His	Gln	Val	Leu	Arg	Trp	Ala	Leu	Gly	Ser	Val	Glu	Ile	Phe	
			820					825					830			
Leu	Ser	Arg	His	Cys	Pro	Ile	Trp	Tyr	Gly	Tyr	Gly	Gly	Gly	Leu	Lys	
		835					840					845				
Trp	Leu	Glu	Arg	Leu	Ser	Tyr	Ile	Asn	Ser	Val	Val	Tyr	Pro	Trp	Thr	
	850					855					860					
Ser	Leu	Pro	Leu	Ile	Val	Tyr	Cys	Ser	Leu	Pro	Ala	Ile	Cys	Leu	Leu	
865					870					875					880	
Thr	Gly	Lys	Phe	Ile	Val	Pro	Glu	Ile	Ser	Asn	Tyr	Ala	Ser	Ile	Leu	
				885					890					895		
Phe	Met	Ala	Leu	Phe	Ser	Ser	Ile	Ala	Ile	Thr	Gly	Ile	Leu	Glu	Met	
			900					905					910			
Gln	Trp	Gly	Lys	Val	Gly	Ile	Asp	Asp	Trp	Trp	Arg	Asn	Glu	Gln	Phe	
		915					920					925				

Trp Val Ile Gly Gly Val Ser Ala His Leu Phe Ala Leu Phe Gln Gly
 930 935 940
 Leu Leu Lys Val Leu Ala Gly Val Asp Thr Asn Phe Thr Val Thr Ser
 945 950 955 960
 Lys Ala Ala Asp Asp Gly Glu Phe Ser Asp Leu Tyr Leu Phe Lys Trp
 965 970 975
 Thr Ser Leu Leu Ile Pro Pro Met Thr Leu Leu Ile Ile Asn Val Ile
 980 985 990
 Gly Val Ile Val Gly Val Ser Asp Ala Ile Ser Asn Gly Tyr Asp Ser
 995 1000 1005
 Trp Gly Pro Leu Phe Gly Arg Leu Phe Phe Ala Leu Trp Val Ile Ile
 1010 1015 1020
 His Leu Tyr Pro Phe Leu Lys Gly Leu Leu Gly Lys Gln Asp Arg Met
 1025 1030 1035 1040
 Pro Thr Ile Ile Val Val Trp Ser Ile Leu Leu Ala Ser Ile Leu Thr
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 Leu Leu Trp Val Arg Val Asn Pro Phe Val Ala Lys Gly Gly Pro Ile
 1060 1065 1070
 Leu Glu Ile Cys Gly Leu Asp Cys Leu
 1075 1080